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Input Set : A:\406.app.txt

Output Set: N:\CRF3\07092001\I543407.raw

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Output Set: N:\CRF3\07092001\I543407.raw

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			~	NCE:		77 - 1	n 1 -	77.	T 1 -	n 7 -	71 -	T 1.	T7 - 3	Db -	0	C1
		гуs	Leu	Leu	_	vaı	Ата	Ата	TTE		Ата	IIe	vaı	Pne		GIA
143	1	70.1 -	T	n 1 -	5	77 - 1	T7 - 1	D	C1-	10	C1	C1	C1	C1	15	114 -
	Ser	Ата	ьeu	Ala	GTÀ	vaı	vai	Pro		Tyr	сту	GIY	GIA	_	ASI	HIS
145	C1	C1	C1	20	7	7	C	C1	25	700	Com	C1	T 0	30	т1.	т
	GTĀ	СТУ	_	Gly	ASII	ASII	Set	40	PIO	ASII	ser	GLU	45	MSII.	шe	ıyı
147 35 40 45 148 Gln Tyr Gly Gly Gly Asn Ser Ala Leu Ala Leu Gln Thr Asp										7 l s	Ara					
149	GIII	50	GTÅ	СТУ	дту	no II	55	лта	пец	лта	пеп	60	1111	тэр	Ата	ALG
	Δsn		Δsn	Leu	Thr	Tle		Gln	His	Glv	Glv		Asn	Glv	Δla	Asp
151		DCI	7150	БСС	1111	70	1111	0111		O _T y	75	O-y	11011	017		80
		Glv	Gln	Gly	Ser		Asp	Ser	Ser	Ile		Leu	Thr	Gln	Ara	
153		0 -1		1	85					90					95	2
	Phe	Glv	Asn	Ser		Thr	Leu	asp	Gln	Trp	Asn	Gly	Lys	Asn	Ser	Glu
155		- 1		100				•	105	•		-	•	110		
	Met	Thr	Val	Lys	Gln	Phe	Gly	Gly	Gly	Asn	Gly	Ala	Ala	Val	Asp	Gln
157			115	-			-	120^{-}	-		-		125		-	
	Thr	Ala	Ser	Asn	Ser	Ser	Val	Asn	Val	Thr	Gln	Val	Gly	Phe	Gly	Asn
159		130					135					140			_	
			Thr	Ala	His	Gln	Tyr									
	145					150										
163	<210	0> SI	EQ II	ON C	: 8											
164	<21	1> L!	ENGTI	H: 15												
165	<21	2> T	YPE:	PRT												
166	<21	3> 01	RGAN	ISM:	: Escherichia co				li							
				NCE:			•									
169	Met	Lys	Asn	Lys	Leu	Leu	Phe	Met	Met	Leu	Thr	Ile	Leu	Gly	Ala	Pro
170	1				5					10					15	

Input Set : A:\406.app.txt

Output Set: N:\CRF3\07092001\I543407.raw

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171 Gly Ile Ala Ala Ala Gly Tyr Asp Leu Ala Asn Ser Glu Tyr Asn
172
173 Phe Ala Val Asn Glu Leu Ser Lys Ser Ser Phe Asn Gln Ala Ala Ile
174
            35
                                40
175 Ile Gly Gln Ala Gly Thr Asn Asn Ser Ala Gln Leu Arg Gln Gly Gly
                            55
177 Ser Lys Leu Leu Ala Val Val Ala Gln Glu Gly Ser Ser Asn Arg Ala
                        70
                                             75
179 Lys Ile Asp Gln Thr Gly Asp Tyr Asn Leu Ala Tyr Ile Asp Gln Ala
                    85
                                        90
181 Gly Ser Ala Asn Asp Ala Ser Ile Ser Gln Gly Ala Tyr Gly Asn Thr
                                    105
183 Ala Met Ile Ile Gln Lys Gly Ser Gly Asn Lys Ala Asn Ile Thr Gln
184
                                120
185 Tyr Gly Thr Gln Lys Thr Ala Ile Val Val Gln Arg Gln Ser Gln Met
                            135
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187 Ala Ile Arg Val Thr Gln Arg
190 <210> SEQ ID NO: 9
191 <211> LENGTH: 48
192 <212> TYPE: DNA
193 <213> ORGANISM: Leishmania major
195 <400> SEQUENCE: 9
196 tatgatcagc tggttacccg tgttgttacc catgaaatgg cacatgca
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198 <210> SEO ID NO: 10
199 <211> LENGTH: 16
200 <212> TYPE: PRT
201 <213> ORGANISM: Leishmania major
203 <400> SEQUENCE: 10
204 Tyr Asp Gln Leu Val Thr Arg Val Val Thr His Glu Met Ala His Ala
205 1
                     5
                                        10
207 <210> SEQ ID NO: 11
208 <211> LENGTH: 456
209 <212> TYPE: DNA
210 <213> ORGANISM: Artificial Sequence
212 <220> FEATURE:
213 <223> OTHER INFORMATION: Recombinant Salmonella enteritidis 3b afgA
214
          sequence containing the replacement fragment
          encoding PT3 from GP63 of Leishmania major.
217 <400> SEQUENCE: 11
218 atgaaacttt taaaagtggc agcattcgca gcaatcgtag tttctggcag tgctctggct
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219 ggcgtcgttc cacaatgggg cggcggcgt aatcataacg gcggcggcaa tagttccggc
                                                                           120
220 ccggactcaa cgttgagcat ttatcagtac ggttccgcta acgctgcgct tgctctgcaa
                                                                           180
221 agegatgeec gtaaatetga aacgaceatt acceagageg gttatggtaa eggegeegat
                                                                           240
222 gtaggccagg gtgcggataa tagtactatt gaactgactc agaatggttt cagaaataat
                                                                           300
223 gccaccatcg accagtggaa cgctaaaaac tccgatatta ctgtcggcca atacggcggt
                                                                           360
224 aataacgccg cgctggttaa ttatgatcag ctggttaccc gtgttgttac ccatgaaatg
                                                                           420
225 gcacatgcaa acaacgccac ggctaaccag tattaa
                                                                           456
227 <210> SEQ ID NO: 12
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Input Set : A:\406.app.txt

Output Set: N:\CRF3\07092001\I543407.raw

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228 <211> LENGTH: 151
229 <212> TYPE: PRT
230 <213> ORGANISM: Artificial Sequence
232 <220> FEATURE:
233 <223> OTHER INFORMATION: Recombinant Salmonella enteritidis 3b afgA
          sequence containing the replacement fragment
234
          encoding PT3 from GP63 of Leishmania major.
235
237 <400> SEQUENCE: 12
238 Met Lys Leu Leu Lys Val Ala Ala Phe Ala Ala Ile Val Val Ser Gly
                                         10
240 Ser Ala Leu Ala Gly Val Val Pro Gln Trp Gly Gly Gly Asn His
242 Asn Gly Gly Gly Asn Ser Ser Gly Pro Asp Ser Thr Leu Ser Ile Tyr
243
244 Gln Tyr Gly Ser Ala Asn Ala Ala Leu Ala Leu Gln Ser Asp Ala Arg
                            55
246 Lys Ser Glu Thr Thr Ile Thr Gln Ser Gly Tyr Gly Asn Gly Ala Asp
                        70
248 Val Gly Gln Gly Ala Asp Asn Ser Thr Ile Glu Leu Thr Gln Asn Gly
                    85
                                         90
250 Phe Arg Asn Asn Ala Thr Ile Asp Gln Trp Asn Ala Lys Asn Ser Asp
                100
                                     105
252 Ile Thr Val Gly Gln Tyr Gly Gly Asn Asn Ala Ala Leu Val Asn Tyr
                                 120
254 Asp Gln Leu Val Thr Arg Val Val Thr His Glu Met Ala His Ala Asn
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                            135
256 Asn Ala Thr Ala Asn Gln Tyr
257 145
                        150
259 <210> SEQ ID NO: 13
260 <211> LENGTH: 456
261 <212> TYPE: DNA
262 <213> ORGANISM: Artificial Sequence
264 <220> FEATURE:
265 <223> OTHER INFORMATION: Recombinant Salmonella enteritidis 3b afgA
          sequence containing the replacement fragment
266
          encoding PT3 from GP63 of Leishmania major...
267
269 <400> SEQUENCE: 13
270 atqaaacttt taaaagtggc agcattcgca gcaatcgtag tttctggcag tgctctggct
                                                                            60
271 ggcgtcgttc cacaatgggg cggcggcggt aatcataacg gcggcggcaa tagttccggc
                                                                           120
272 ccqqactcaa cqttqaqcat ttatcaqtac ggttccqcta acgctqcqct tqctctqcaa
                                                                           180
273 agcqatqccc qtaaatctqa aacqaccatt acccagagcg gttatggtaa cggcgccgat
                                                                           240
274 gtaggccagg gtgcggataa tagtactatt gaactgactc agaatggttt cagaaataat
                                                                           300
275 gccaccatcg accagtggaa cgctaaaaac tccgatatta ctgtcggcca atatgatcag
                                                                           360
                                                                           420
276 ctqqttaccc gtgttgttac ccatgaaatg gcacatgcaa gcgtaatggt gcgtcaggtt
                                                                           456
277 ggttttggca acaacgccac ggctaaccag tattaa
279 <210> SEQ ID NO: 14
280 <211> LENGTH: 151
281 <212> TYPE: PRT
282 <213> ORGANISM: Artificial Sequence
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Use of n and/or Xaa has been detected in the Sequence Listing.

Review the Sequence Listing to insure a conesponding explanation is presented in the <220> to <223> fields of each sequence using n or Xaa.

VERIFICATION SUMMARY

DATE: 07/09/2001 TIME: 12:43:43

PATENT APPLICATION: US/09/543,407

Input Set : A:\406.app.txt

Output Set: N:\CRF3\07092001\I543407.raw

L:783 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33 L:785 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33 L:1147 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:59 L:1149 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:59